AMENDMENT

The following <u>Listing</u> of the <u>Claims</u> will replace all prior versions and all prior listings of the claims in the present application:

Listing of The Claims:

Claim 1 – 19 (Cancelled)

Claim 20 (Currently Amended): A method for diagnosing the risk of restenosis after coronary angioplasty in a Japanese man, comprising the following steps (i) to (iii):

- (i) analyzing the following polymorphisms (1), (3) and (4) in a human nucleic acid sample from said Japanese man;
- (1) polymorphism at the base number position 3932 (the 3932nd base of SEQ ID NO:1) of the human apolipoprotein E gene;
- (3) polymorphism at the base number position -863 (the 197th base of SEQ ID NO:3)of the human tumor necrosis factor- α gene;
- (4) polymorphism at the base number position 825 (the 831st base of SEQ ID NO:4)of the human G-protein β 3 subunit gene;
- (ii) determining, based on the information about polymorphism which was obtained in the step (i), the genotype of the nucleic acid sample; and
- (iii) assessing, based on the genotype determined, a genetic risk of restenosis after coronary angioplasty in a Japanese man.